



SEQUENCE LISTING

<110> Pepinsky, et al.

<120> HYDROPHOBICALLY-MODIFIED PROTEIN COMPOSITIONS AND METHODS

<130> BIIJ-P02-067

<140> 09/579,680

<141> 2000-05-26

<150> PCT/US98/25676

<151> 1998-12-13

<160> 4

<170> PatentIn Ver. 3.1

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<213> human

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20 25 30Thr Leu Gly Ala Ser Gly Arg Tyr Glu Gly Lys Ile Ala Arg Ser Ser
35 40 45Glu Arg Phe Lys Glu Leu Thr Pro Asn Tyr Asn Pro Asp Ile Ile Phe
50 55 60Lys Asp Glu Glu Asn Thr Gly Ala Asp Arg Leu Met Thr Gln Arg Cys
65 70 75 80Lys Asp Arg Leu Asn Ser Leu Ala Ile Ser Val Met Asn Gln Trp Pro
85 90 95Gly Val Lys Leu Arg Val Thr Glu Gly Trp Asp Glu Asp Gly His His
100 105 110Ser Glu Glu Ser Leu His Tyr Glu Gly Arg Ala Val Asp Ile Thr Thr
115 120 125Ser Asp Arg Asp Arg Asn Lys Tyr Gly Leu Leu Ala Arg Leu Ala Val
130 135 140Glu Ala Gly Phe Asp Trp Val Tyr Tyr Glu Ser Lys Ala His Val His
145 150 155 160Cys Ser Val Lys Ser Glu His Ser Ala Ala Ala Lys Thr Gly Gly
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35 40 45
Arg Phe Lys Glu Leu Thr Pro Asn Tyr Asn Pro Asp Ile Ile Phe Lys
50 55 60
Asp Glu Glu Asn Thr Gly Ala Asp Arg Leu Met Thr Gln Arg Cys Lys
65 70 75 80
Asp Lys Leu Asn Ala Leu Ala Ile Ser Val Met Asn Gln Trp Pro Gly
85 90 95
Val Lys Leu Arg Val Thr Glu Gly Trp Asp Glu Asp Gly His His Ser
100 105 110
Glu Glu Ser Leu His Tyr Glu Gly Arg Ala Val Asp Ile Thr Thr Ser
115 120 125
Asp Arg Asp Arg Ser Lys Tyr Gly Met Leu Ala Arg Leu Ala Val Glu
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Ala Gly Phe Asp Trp Val Tyr Tyr Glu Ser Lys Ala His Ile His Cys
145 150 155 160
Ser Val Lys Ala Glu Asn Ser Val Ala Ala Lys Ser Gly Gly
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Arg Thr Leu Gly Ala Ser Gly Pro Ala Glu Gly Arg Val Ala Arg Gly
35 40 45

Ser	Glu	Arg	Phe	Arg	Asp	Leu	Val	Pro	Asn	Tyr	Asn	Pro	Asp	Ile	Ile
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Phe	Lys	Asp	Glu	Glu	Asn	Ser	Gly	Ala	Asp	Arg	Leu	Met	Thr	Glu	Arg
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Cys	Lys	Glu	Arg	Val	Asn	Ala	Leu	Ala	Ile	Ala	Val	Met	Asn	Met	Trp
				85					90				95		
Pro	Gly	Val	Arg	Leu	Arg	Val	Thr	Glu	Gly	Trp	Asp	Glu	Asp	Gly	His
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His	Ala	Gln	Asp	Ser	Leu	His	Tyr	Glu	Gly	Arg	Ala	Leu	Asp	Ile	Thr
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Thr	Ser	Asp	Arg	Asp	Arg	Asn	Lys	Tyr	Gly	Leu	Leu	Ala	Arg	Leu	Ala
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Val	Glu	Ala	Gly	Phe	Asp	Trp	Val	Tyr	Tyr	Glu	Ser	Arg	Asn	His	Val
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Xaa Leu Xaa Pro Leu Xaa Tyr Lys Gln Phe Xaa Pro Xaa Val Xaa Glu
20 25 30

Lys Thr Leu Gly Ala Ser Gly Arg Xaa Glu Gly Lys Xaa Xaa Arg Xaa
35 40 45

Ser Glu Arg Phe Lys Xaa Leu Xaa Pro Asn Tyr Asn Pro Asp Ile Ile
50 55 60

Phe Lys Asp Glu Glu Asn Xaa Gly Ala Asp Arg Leu Met Thr Xaa Arg
65 70 75 80

Cys Lys Xaa Xaa Xaa Asn Ser Leu Ala Ile Xaa Val Met Asn Xaa Trp
85 90 95

Pro Gly Val Lys Leu Arg Val Thr Glu Gly Trp Asp Glu Asp Gly His
100 105 110

His Xaa Xaa Xaa Ser Leu His Tyr Glu Gly Arg Ala Val Asp Ile Thr
115 120 125

Thr Ser Asp Arg Asp Arg Xaa Lys Tyr Gly Xaa Leu Ala Arg Leu Ala
130 135 140

Val Glu Ala Gly Phe Asp Trp Val Tyr Tyr Glu Ser Xaa Xaa His Xaa
145 150 155 160

His Xaa Ser Val Lys Xaa Xaa Xaa Ser Xaa Ala Ala Xaa Xaa Gly Gly
165 170 175

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